This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims: Please amend the claims as follows:

We claim:

Claims 1.-53. (Cancelled)

Claim 54. (Currently Amended) A process for the production of ergosterol or an intermediate product thereof which is squalene, farnesol, geraniol, lanosterol, zymosterol, 4,4dimethylzymosterol, 4-methylzymosterol, ergost-7-enol or ergosta-5,7-dienol or a sterol with a 5,7-diene structure, comprising fermenting into ergosterol with a plasmid vector-transformed microorganism, wherein said vector comprises several suitable genes of the ergosterol metabolic process in altered form, wherein the catalytic area of HMG is expressed without its membranebonded domain; and the natural promoter of the gene t-HMG, ERG9 and SAT1 gene is replaced by the middle part of ADH1 promoter, the suitable genes being

a-i) i) the a gene of the HMG-Co-A-reductase (t-HMG), i) ii) the a gene of the squalene synthetase (ERG9), ii) iii) the a gene of the acyl-CoA; sterol-acyltransferase (SAT1), and iii) iv) the a gene of squalene epoxidase (ERG1), or a-ii) i) the a gene of HMG-Co-A-reductase (t-HMG), and ii) the a gene of the squalene synthetase (ERG9), or a-iii) i) the a gene of the HMG-Co-A-reductase (t-HMG), and ii) the a gene of the acyl-CoA: sterol-acyltransferase (SAT1), or a-iv)

ii) the a gene of squalene epoxidase (ERG1),

or

i)

the a gene of the HMG-Co-A-reductase (t-HMG), and

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a-v)
               the a gene of squalene synthetase (ERG9), and
       i)
               the a gene of acyl-CoA: sterol-acyltransferase (SAT1)
        ii)
or
a-vi)
               the a gene of squalene synthetase (ERG9), and
        i)
               the a gene of squalene epoxidase (ERG1),
        ii)
or
a-vii)
               the a gene of acyl-coA: sterol-acyltransferase (SAT1) and
        iii) i)
        i) ii)
               the a gene of squalene epoxidase (ERG1),
or
a-viii) at least one of the genes of a-i),
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analyzing, purifying and isolating the fermented product.

Claim 55. (Currently Amended) A process according to claim 54, further comprising the <u>a</u> gene of squalene epoxidase (*ERG1*) in a-ii), a-iii), and a-v), and the <u>a</u> gene of acyl-CoA: sterol-acyl-transferase in a-ii).

Claim 56. (Previously Presented) A process according to claim 54, wherein the genes in a-i) to a-vii) are introduced, in each case independently of one another, into microorganisms of the same species.

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Claim 57. (Cancelled)
Claim 58. (Cancelled)
Claim 59. (Cancelled)
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and

Claim 60. (Previously Presented) A process according to claim 54, wherein the microorganism is yeast.

Claim 61. (Previously Presented) A process according to claim 60, wherein said yeast is S. cerevisiae.

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Claim 62. (Previously Presented) A process according to claim 61, wherein said yeast is S. cerevisiae AH22.

Claim 63. (Previously Presented) Yeast strain S. cerevisiae AH22, comprising one or more of the genes mentioned under a-i) of claim 54.

Claim 64. (Cancelled)

Claim 65. (Cancelled)

Claim 66. (Cancelled)

Claim 67. (Cancelled)

Claim 68. (Previously Presented) An expression cassette that comprises the average ADH promoter, the t-HMG gene, the TRP terminator and the SAT1 gene with the average ADH promoter and the TRP terminator.

Claim 69. (Previously Presented) An expression cassette comprising the average ADH promoter, the t-HMG gene, the TRP terminator, the SAT1 gene with the average ADH promoter, and the TRP terminator, and the ERG9 gene with the average ADH promoter and the TRP terminator.

Claim 70. (Previously Presented) A combination of expression cassettes comprising

- a) a first expression cassette, on which the *ADH* promoter, the *t-HMG* gene and the *TRP* terminator are located,
- b) a second expression cassette, on which the *ADH* promoter, the *SAT1* gene and the *TRP* terminator are located,

and

a third expression cassette, on which the *ADH* promoter, the *ERG9* gene with the *TRP* terminator are located.

Claim 71. (Cancelled)

Claim 72. (Cancelled)

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Claim 73. (Previously Presented) A microorganism comprising an expression cassette according to claim 68.

Claim 74. (Previously Presented) A microorganism according to claim 73, which is yeast.

Claim 75. (Cancelled)

Claim 76. (Cancelled)

Claim 77. (Previously Presented) A process according to claim 55, wherein the genes are introduced, in each case independently of one another, into microorganisms of the same species.

Claim 78. (Previously Presented) A microorganism comprising an expression cassette according to claim 69.

Claim 79. (Previously Presented) A microorganism comprising a combination according to claim 70.

Claim 80. (Previously Presented) A microorganism according to claim 78, which is yeast.

Claim 81. (Previously Presented) A microorganism according to claim 79, which is yeast.

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